RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/575,537_

Source: _TEMP_

Date Processed by STIC: _9/21/06_

ENTERED

RAW SEQUENCE LISTING   DATE: 09/21/2006
PATENT APPLICATION: US/10/575,537   TIME: 11:48:13

Input Set : A:\1034123-000219.ST25.txt
Output Set: N:\CRF4\09212006\J575537.raw

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      3 <110> APPLICANT: Gallo, Richard
      4          Murakami, Masamoto
      6 <120> TITLE OF INVENTION: HUMAN CATHELICIDIN ANTIMICROBIAL PEPTIDES
      8 <130> FILE REFERENCE: 1034123-000219
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/575,537
     11 <141> CURRENT FILING DATE: 2006-04-11
     13 <150> PRIOR APPLICATION NUMBER: US 60/512,953
     14 <151> PRIOR FILING DATE: 2003-10-21
     16 <150> PRIOR APPLICATION NUMBER: PCT/US2004/034911
     17 <151> PRIOR FILING DATE: 2004-10-20
     19 <160> NUMBER OF SEQ ID NOS: 34
     21 <170> SOFTWARE: PatentIn version 3.3
     23 <210> SEQ ID NO: 1
     24 <211> LENGTH: 11
     25 <212> TYPE: PRT
     26 <213> ORGANISM: Homo sapiens
     29 <220> FEATURE:
W--> 30 <221> NAME/KEY: X
     31 <222> LOCATION: (1)..(2)
     32 <223> OTHER INFORMATION: K or R
     34 <220> FEATURE:
W--> 35 <221> NAME/KEY: X
     36 <222> LOCATION: (3)..(3)
     37 <223> OTHER INFORMATION: I or K
     39 <220> FEATURE:
W--> 40 <221> NAME/KEY: X
     41 <222> LOCATION: (4)..(4)
     42 <223> OTHER INFORMATION: V or G
     44 <220> FEATURE:
W--> 45 <221> NAME/KEY: X
     46 <222> LOCATION: (5)..(5)
     47 <223> OTHER INFORMATION: Q or R
     49 <220> FEATURE:
W--> 50 <221> NAME/KEY: X
     51 <222> LOCATION: (6)..(6)
     52 <223> OTHER INFORMATION: K or R
     54 <220> FEATURE:
W--> 55 <221> NAME/KEY: X
     56 <222> LOCATION: (7)..(7)
     57 <223> OTHER INFORMATION: any amino acid
     59 <220> FEATURE:
W--> 60 <221> NAME/KEY: X
     61 <222> LOCATION: (8)..(8)
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       64 <220> FEATURE:
W-->   65 <221> NAME/KEY: X
       66 <222> LOCATION: (9)..(11)
       67 <223> OTHER INFORMATION: any amino acid
       69 <400> SEQUENCE: 1
W-->   71 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
       72 1               5                   10
       75 <210> SEQ ID NO: 2
       76 <211> LENGTH: 11
       77 <212> TYPE: PRT
       78 <213> ORGANISM: Homo sapiens
       80 <400> SEQUENCE: 2
       82 Lys Arg Ile Val Gln Arg Ile Lys Asp Val Phe
       83 1               5                   10
       86 <210> SEQ ID NO: 3
       87 <211> LENGTH: 8
       88 <212> TYPE: PRT
       89 <213> ORGANISM: Homo sapiens
       91 <400> SEQUENCE: 3
       93 Arg Lys Ser Lys Glu Lys Ile Gly
       94 1               5
       97 <210> SEQ ID NO: 4
       98 <211> LENGTH: 8
       99 <212> TYPE: PRT
      100 <213> ORGANISM: Homo sapiens
      102 <400> SEQUENCE: 4
      104 Lys Ser Lys Glu Lys Ile Gly Lys
      105 1               5
      108 <210> SEQ ID NO: 5
      109 <211> LENGTH: 739
      110 <212> TYPE: DNA
      111 <213> ORGANISM: Homo sapiens
      113 <400> SEQUENCE: 5
      114 taaagcaaac cccagcccac accctggcag gcagccaggg atgggtggat caggaaggct      60
      116 cctggttggg cttttgcatc aggctcaggc tgggcataaa ggaggctcct gtgggctaga     120
      118 gggaggcaga catgggacc atgaagaccc aaagggatgg ccactccctg gggcggtggt     180
      120 cactggtgct cctgctgctg ggcctggtga tgcctctggc catcattgcc caggtcctca     240
      122 gctacaagga agctgtgctt cgtgctatag atggcatcaa ccagcggtcc tcggatgcta     300
      124 acctctaccg cctcctggac ctggacccca ggcccacgat ggatggggac ccagacacgc     360
      126 caaagcctgt gagcttcaca gtgaaggaga cagtgtgccc caggacgaca cagcagtcac     420
      128 cagaggattg tgacttcaag aaggacgggc tggtgaagcg gtgtatgggg acagtgaccc     480
      130 tcaaccaggc caggggctcc tttgacatca gttgtgataa ggataacaag agatttgccc     540
      132 tgctgggtga tttcttccgg aaatctaaag agaagattgg caaagagttt aaaagaattg     600
      134 tccagagaat caaggatttt ttgcggaatc ttgtacccag acagagtcc tagtgtgtgc     660
      136 cctaccctgg ctcaggcttc tgggctctga gaaataaact atgagagcaa tttcaaaaaa     720
      138 aaaaaaaaaa aaaaaaaaa                                                  739
      141 <210> SEQ ID NO: 6
      142 <211> LENGTH: 170
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Input Set : A:\1034123-000219.ST25.txt
Output Set: N:\CRF4\09212006\J575537.raw

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144 <213> ORGANISM: Homo sapiens
146 <400> SEQUENCE: 6
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152 Leu Leu Leu Leu Gly Leu Val Met Pro Leu Ala Ile Ile Ala Gln Val
153             20                  25                  30
156 Leu Ser Tyr Lys Glu Ala Val Leu Arg Ala Ile Asp Gly Ile Asn Gln
157         35                  40                  45
160 Arg Ser Ser Asp Ala Asn Leu Tyr Arg Leu Leu Asp Leu Asp Pro Arg
161     50                  55                  60
164 Pro Thr Met Asp Gly Asp Pro Asp Thr Pro Lys Pro Val Ser Phe Thr
165 65                  70                  75                  80
168 Val Lys Glu Thr Val Cys Pro Arg Thr Thr Gln Gln Ser Pro Glu Asp
169                 85                  90                  95
172 Cys Asp Phe Lys Lys Asp Gly Leu Val Lys Arg Cys Met Gly Thr Val
173             100                 105                 110
176 Thr Leu Asn Gln Ala Arg Gly Ser Phe Asp Ile Ser Cys Asp Lys Asp
177         115                 120                 125
180 Asn Lys Arg Phe Ala Leu Leu Gly Asp Phe Phe Arg Lys Ser Lys Glu
181     130                 135                 140
184 Lys Ile Gly Lys Glu Phe Lys Arg Ile Val Gln Arg Ile Asp Asp Phe
185 145                 150                 155                 160
188 Leu Arg Asn Leu Val Pro Arg Thr Glu Ser
189                 165                 170
192 <210> SEQ ID NO: 7
193 <211> LENGTH: 519
194 <212> TYPE: DNA
195 <213> ORGANISM: Homo sapiens
197 <400> SEQUENCE: 7
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200 ctactgggcc tggggttctc ccagaccccc agctacaggg atgctgtgct ccgagctgtg     120
202 gatgacttca ccagcagtc  cctagacacc aatctctacc gtctcctgga cctggatcct     180
204 gagccccaag gggacgagga tccagatact cccaagtctg tgaggttccg agtgaaggag     240
206 actgtatgtg gcaaggcaga gcggcagcta cctgagcaat gtgccttcaa ggaacagggg     300
208 gtggtgaagc agtgtatggg ggcagtcacc ctgaacccgg ccgctgattc ttttgacatc     360
210 agctgtaacg agcctggtgc acagcccttt cggttcaaga aaatttcccg gctggctgga     420
212 cttctccgca aggtggggga aagattggt  gaaaagctta agaaaattgg ccagaaaatt     480
214 aagaattttt ttcagaaact tgtccctcag ccagagtag                            519
217 <210> SEQ ID NO: 8
218 <211> LENGTH: 173
219 <212> TYPE: PRT
220 <213> ORGANISM: murine
222 <400> SEQUENCE: 8
224 Met Gln Phe Gln Arg Asp Val Pro Ser Leu Trp Leu Trp Arg Ser Leu
225 1               5                   10                  15
228 Ser Leu Leu Leu Leu Leu Gly Leu Gly Phe Ser Gln Thr Pro Ser Tyr
229             20                  25                  30
232 Arg Asp Ala Val Leu Arg Ala Val Asp Asp Phe Asn Gln Gln Ser Leu
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```
233         35                    40                    45
236 Asp Thr Asn Leu Tyr Arg Leu Leu Asp Leu Asp Pro Glu Pro Gln Gly
237    50                    55                    60
240 Asp Glu Asp Pro Asp Thr Pro Lys Ser Val Arg Phe Arg Val Lys Glu
241 65                    70                    75                    80
244 Thr Val Cys Gly Lys Ala Glu Arg Gln Leu Pro Glu Gln Cys Ala Phe
245                   85              90                    95
248 Lys Glu Gln Gly Val Val Lys Gln Cys Met Gly Ala Val Thr Leu Asn
249                   100             105                   110
252 Pro Ala Ala Asp Ser Phe Asp Ile Ser Cys Asn Glu Pro Gly Ala Gln
253         115                   120                   125
256 Pro Phe Arg Phe Lys Lys Ile Ser Arg Leu Ala Gly Leu Leu Arg Lys
257    130                   135                   140
260 Gly Gly Glu Lys Ile Gly Glu Lys Leu Lys Lys Ile Gly Gln Lys Ile
261 145                   150                   155                   160
264 Lys Asn Phe Phe Gln Lys Leu Val Pro Gln Pro Glu Gln
265                   165                   170
268 <210> SEQ ID NO: 9
269 <211> LENGTH: 172
270 <212> TYPE: PRT
271 <213> ORGANISM: canine
273 <400> SEQUENCE: 9
275 Met Glu Thr Gln Lys Asp Ser Pro Ser Leu Gly Arg Trp Ser Leu Leu
276 1                   5                   10                    15
279 Leu Leu Leu Leu Gly Leu Val Ile Thr Pro Ala Ala Ser Arg Ala Leu
280                   20                    25                    30
283 Ser Tyr Arg Glu Ala Val Leu Arg Ala Val Asn Gly Phe Asn Gln Arg
284         35                    40                    45
287 Ser Ser Glu Glu Asn Leu Tyr Arg Leu Leu Gln Leu Asn Ser Gln Pro
288    50                    55                    60
291 Lys Gly Asp Glu Asp Pro Asn Ile Pro Lys Pro Val Ser Phe Thr Val
292 65                    70                    75                    80
295 Lys Glu Thr Val Cys Pro Lys Thr Thr Gln Gln Pro Leu Glu Gln Cys
296                   85                    90                    95
299 Gly Phe Lys Asp Asn Gly Leu Val Lys Gln Cys Glu Gly Thr Val Ile
300                   100                   105                   110
303 Leu Asp Glu Asp Thr Gly Tyr Phe Asp Leu Asn Cys Asp Ser Ile Leu
304         115                   120                   125
307 Gln Val Lys Lys Ile Asp Arg Leu Lys Glu Leu Ile Thr Thr Gly Ala
308    130                   135                   140
311 Gln Lys Ile Gly Lys Lys Ile Arg Arg Ile Gly Gln Arg Ile Lys Asp
312 145                   150                   155                   160
315 Phe Leu Lys Asn Leu Gln Pro Arg Glu Glu Lys Ser
316                   165                   170
319 <210> SEQ ID NO: 10
320 <211> LENGTH: 172
321 <212> TYPE: PRT
322 <213> ORGANISM: porcine
324 <400> SEQUENCE: 10
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Input Set : A:\1034123-000219.ST25.txt
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```
326 Met Glu Thr Gln Arg Ala Ser Leu Cys Leu Gly Arg Trp Ser Leu Trp
327 1               5                  10                  15
330 Leu Leu Leu Leu Ala Leu Val Val Pro Ser Ala Ser Ala Gln Ala Leu
331             20                  25                  30
334 Ser Tyr Arg Glu Ala Val Leu Arg Ala Val Asp Arg Leu Asn Glu Gln
335         35                  40                  45
338 Ser Ser Glu Ala Asn Leu Tyr Arg Leu Leu Glu Leu Asp Gln Pro Pro
339      50                  55                  60
342 Lys Ala Asp Glu Asp Pro Gly Thr Pro Lys Pro Val Ser Phe Thr Val
343 65                  70                  75                  80
346 Lys Glu Thr Val Cys Pro Arg Pro Thr Arg Gln Pro Pro Glu Leu Cys
347             85                  90                  95
350 Asp Phe Lys Glu Asn Gly Arg Val Lys Gln Cys Val Gly Thr Val Thr
351         100                 105                 110
354 Leu Asn Pro Ser Ile His Ser Leu Asp Ile Ser Cys Asn Glu Ile Gln
355         115                 120                 125
358 Ser Val Arg Arg Arg Pro Arg Pro Pro Tyr Leu Pro Arg Pro Arg Pro
359     130                 135                 140
362 Pro Pro Phe Phe Pro Pro Arg Leu Pro Pro Arg Ile Pro Pro Gly Phe
363 145                 150                 155                 160
366 Pro Pro Arg Phe Pro Pro Arg Phe Pro Gly Lys Arg
367             165                 170
370 <210> SEQ ID NO: 11
371 <211> LENGTH: 176
372 <212> TYPE: PRT
373 <213> ORGANISM: goat
375 <400> SEQUENCE: 11
377 Met Glu Thr Gln Gly Ala Ser Leu Ser Leu Gly Arg Trp Ser Leu Trp
378 1               5                  10                  15
381 Leu Leu Leu Leu Gly Leu Val Val Pro Leu Ala Ser Ala Gln Ala Leu
382             20                  25                  30
385 Ser Tyr Arg Glu Ala Val Leu Arg Ala Val Gly Gln Leu Asn Glu Arg
386         35                  40                  45
389 Ser Ser Glu Ala Asn Leu Tyr Arg Leu Leu Glu Leu Asp Pro Ala Pro
390     50                  55                  60
393 Asn Asp Glu Val Asp Pro Gly Thr Arg Lys Pro Val Ser Phe Thr Val
394 65                  70                  75                  80
397 Lys Glu Thr Val Cys Pro Arg Thr Thr Gln Gln Pro Pro Glu Glu Cys
398             85                  90                  95
401 Asp Phe Lys Glu Asn Gly Leu Val Lys Gln Cys Val Gly Thr Val Thr
402         100                 105                 110
405 Leu Asp Pro Ser Asn Asp Gln Phe Asp Ile Asn Cys Asn Glu Leu Gln
406         115                 120                 125
409 Ser Val Arg Phe Arg Pro Pro Ile Arg Arg Pro Pro Ile Arg Pro Pro
410     130                 135                 140
413 Phe Asn Pro Pro Phe Arg Pro Pro Val Arg Pro Pro Phe Arg Pro Pro
414 145                 150                 155                 160
417 Phe Arg Pro Pro Phe Arg Pro Pro Ile Gly Pro Phe Pro Gly Arg Arg
418             165                 170                 175
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 09/21/2006
PATENT APPLICATION: **US/10/575,537**   TIME: 11:48:14

Input Set : **A:\1034123-000219.ST25.txt**
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Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11

VERIFICATION SUMMARY
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Input Set : **A:\1034123-000219.ST25.txt**
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L:10 M:270 C: Current Application Number differs, Replaced Current Application Number
L:30 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:35 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:40 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:45 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:50 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:55 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:60 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:65 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:71 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0